

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

08/894356

Art Unit / Team No. :

1649

Date Processed by STIC:

3/25/98

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

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1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,

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THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

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ARTI SHAH 703-308-4212

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/894,356DATE: 03/25/98
TIME: 11:00:45

INPUT SET: S24372.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: ASHIKARI, Toshihiko
TANAKA, Yoshikazu
FUJIWARA, Hiroyuki
NAKAO, Masahiro
FUKUI, Yuko
SAKAKIBARA, Keiko
MIZUTANI, Masako
KUSUMI, Takaaki

(ii) TITLE OF INVENTION: GENE ENCODING A PROTEIN HAVING ACYL
GROUP TRANSFER ACTIVITY

(iii) NUMBER OF SEQUENCES: 31

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS, L.L.P.
(B) STREET: P.O. Box 1404
(C) CITY: Alexandria
(D) STATE: Virginia
(E) COUNTRY: United States
(F) ZIP: 22313-1404

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US 08/894,356
(B) FILING DATE: 18-AUG-1997
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: JP 7-67159
(B) FILING DATE: 17-FEB-1995

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: JP 7-196915
(B) FILING DATE: 29-JUN-1995

Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/894,356DATE: 03/25/98
TIME: 11:00:47

INPUT SET: S24372.raw

46 (vii) PRIOR APPLICATION DATA:
47 (A) APPLICATION NUMBER: JP 8-46534
48 (B) FILING DATE: 30-JAN-1996
49
50 (vii) PRIOR APPLICATION DATA:
51 (A) APPLICATION NUMBER: WO PCT/JP96/00348
52 (B) FILING DATE: 16-FEB-1996
53
54 (viii) ATTORNEY/AGENT INFORMATION:
55 (A) NAME: Meuth, Donna M.
56 (B) REGISTRATION NUMBER: 36,607
57 (C) REFERENCE/DOCKET NUMBER: 001560-308
58
59 (ix) TELECOMMUNICATION INFORMATION:
60 (A) TELEPHONE: (703) 836-6620
61 (B) TELEFAX: (703) 836-2021
62
63

ERRORED SEQUENCES FOLLOW:

836 (2) INFORMATION FOR SEQ ID NO:6:
837
838 (i) SEQUENCE CHARACTERISTICS:
839 (A) LENGTH: 1521 base pairs
840 (B) TYPE: nucleic acid
841 (C) STRANDEDNESS: double
842 (D) TOPOLOGY: linear
843
844 (ii) MOLECULE TYPE: cDNA to mRNA
845
846 (iii) HYPOTHETICAL: NO
847
848 (iv) ANTI-SENSE: NO
849
850 (vi) ORIGINAL SOURCE:
851 (A) ORGANISM: Lavandula angustifolia
852 (F) TISSUE TYPE: petal
853
854 (vii) IMMEDIATE SOURCE:
855 (A) LIBRARY: cDNA library
856 (B) CLONE: pLAT21
857
858 (ix) FEATURE:
859 (A) NAME/KEY: CDS
860 (B) LOCATION: 3..1352
861
862
863 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
864

next page

RAW SEQUENCE LISTING

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865 -TG ACC ACC CTC CTC GAA TCC TCC CGA GTG GCG CCG CCT CCA GGC ACG
 866 XXX Thr Thr Leu Leu Glu Ser Ser Arg Val Ala Pro Pro Pro Gly Thr
 867 1 1 5 5 10 10 15 15
 868
 869 GTG GCT GAG CAG TCA CTC CCG CTC ACC TTC TTC GAC ATG ACG TGG CTG 95
 870 Val Ala Glu Gln Ser Leu Pro Leu Thr Phe Phe Asp Met Thr Trp Leu
 871 20 25 30
 872
 873 CAT TTC CAC CCC ATG CTT CAG CTT CTC TTC TAC GAA CTC CCC TGT TCC 143
 874 His Phe His Pro Met Leu Gln Leu Phe Tyr Glu Leu Pro Cys Ser
 875 35 40 45
 876
 877 AAA CCC GCC TTC CTC GAA ACC GTC GTT CCG AAA CTC AAA CAA TCC TTA 191
 878 Lys Pro Ala Phe Leu Glu Thr Val Val Pro Lys Leu Lys Gln Ser Leu
 879 50 55 60
 880
 881 TCT CTA ACC CTC AAA CAC TTC TTC CCC CTT TCA TGC AAT CTA ATC TAC 239
 882 Ser Leu Thr Leu Lys His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr
 883 65 70 75
 884
 885 CCT CTA TCG CCG GAG AAA ATG CCG GAG TTC CGG TAT CAG AAC GGT GAC 287
 886 Pro Leu Ser Pro Glu Lys Met Pro Glu Phe Arg Tyr Gln Asn Gly Asp
 887 80 85 90 95
 888
 889 TCG GTT TCT TTC ACG ATT ATG GAG TCT GTC GGA GAT CAT CCG CAT TCC 335
 890 Ser Val Ser Phe Thr Ile Met Glu Ser Val Gly Asp His Pro His Ser
 891 100 105 110
 892
 893 GCT CAT AAA TAC TAC TGC TTT GCC CCT AGC GAC GAT TAT GAA GAT CTC 383
 894 Ala His Lys Tyr Tyr Cys Phe Ala Pro Ser Asp Asp Tyr Glu Asp Leu
 895 115 120 125
 896
 897 CAG CTG CCG CCG ATA GTC GAG GAA TCT GAT CGG AAA TTG TTT CAA GTT 431
 898 Gln Leu Pro Pro Ile Val Glu Glu Ser Asp Arg Lys Leu Phe Gln Val
 899 130 135 140
 900
 901 TTA GCC GTG CAA GTG ACT CTG TTT CCC GGT CGC GGG GTG TGC ATC GGA 479
 902 Leu Ala Val Gln Val Thr Leu Phe Pro Gly Arg Gly Val Cys Ile Gly
 903 145 150 155
 904
 905 ATA ACG ACG CAC CAC ACC GTT AGC GAT GCT CCA TCG TTT GTA GGG TTT 527
 906 Ile Thr Thr His His Thr Val Ser Asp Ala Pro Ser Phe Val Gly Phe
 907 160 165 170 175
 908
 909 ATG AAG AGT TGG GCT TCC ATC ACT AAA TTC GGA GGA GAT GAT GAA TTC 575
 910 Met Lys Ser Trp Ala Ser Ile Thr Lys Phe Gly Gly Asp Asp Glu Phe
 911 180 185 190
 912
 913 TTG GAC GGA AAA GGT GAA TGT TTG CCG GTT TTC GAC CGA TCG CTC GTG 623
 914 Leu Asp Gly Lys Gly Glu Cys Leu Pro Val Phe Asp Arg Ser Leu Val
 915 195 200 205
 916
 917 AAT TAT CCG CCT AAA TTG GAC ACA TAT TTA TGG AAC AAC GCG CAG AAA 671

7 amino numbering starts with the first amino acid

RAW SEQUENCE LISTING PATENT APPLICATION US/08/894,356

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INPUT SET: S24372.raw

918	Asn Tyr Pro Pro Lys Leu Asp Thr Tyr Leu Trp Asn Asn Ala Gln Lys	
919	210 215 220	
920		
921	CGT CCG TTG GAA TCG CAG CAT CCA TCT TTA CCG ACG GAT CGG ATT CGA	719
922	Arg Pro Leu Glu Ser Gln His Pro Ser Leu Pro Thr Asp Arg Ile Arg	
923	225 230 235	
924		
925	GCT ACC TAC CTT TTC ACC CAA TCT GAA ATT AAG AAA TTG AAG GGT TTG	767
926	Ala Thr Tyr Leu Phe Thr Gln Ser Glu Ile Lys Lys Leu Lys Gly Leu	
927	240 245 250 255	
928		
929	ATT CAG AGA AAA GCC CCA AAT GTA GTT AAT CTC TCT TCC TTC GTC GCG	815
930	Ile Gln Arg Lys Ala Pro Asn Val Val Asn Leu Ser Ser Phe Val Ala	
931	260 265 270	
932		
933	ATC GCA GCT TAT ATC TGG ACC GGC ATC GCC AAA TCG GTC GGA GAT TAC	863
934	Ile Ala Ala Tyr Ile Trp Thr Gly Ile Ala Lys Ser Val Gly Asp Tyr	
935	275 280 285	
936		
937	AAA GAC GTG GAT GAC GAC AAA CGC GCT TTC TTT TTA ATT CCG ATC GAT	911
938	Lys Asp Val Asp Asp Asp Lys Arg Ala Phe Phe Leu Ile Pro Ile Asp	
939	290 295 300	
940		
941	TTA AGG CCG CGT TTG GAT CCG CCG GCT CCG GGG AAC TAC TTC GGA AAC	959
942	Leu Arg Pro Arg Leu Asp Pro Pro Ala Pro Gly Asn Tyr Phe Gly Asn	
943	305 310 315	
944		
945	TGT CTA TCG TTT GCG ATG GCG AAG ATC CTG CGG CGG GAT TTG GTC GGA	1007
946	Cys Leu Ser Phe Ala Met Ala Lys Ile Leu Arg Arg Asp Leu Val Gly	
947	320 325 330 335	
948		
949	GAT GAA GGG GTG TTT CGG GCA GCT GAG GCG ATC GCG GCG GAA ATA GAG	1055
950	Asp Glu Gly Val Phe Arg Ala Ala Glu Ala Ile Ala Ala Glu Ile Glu	
951	340 345 350	
952		
953	AAG AGG ACG AGC GAC AAG AAG ATT CTA GAA ACT GTG GAG AAC TGG CCG	1103
954	Lys Arg Thr Ser Asp Lys Lys Ile Leu Glu Thr Val Glu Asn Trp Pro	
955	355 360 365	
956		
957	TCT GAG ATT CGC GAA GCC TTG CAA AAC TGT TAT TTC TCG GTG GCG GGA	1151
958	Ser Glu Ile Arg Glu Ala Leu Gln Asn Cys Tyr Phe Ser Val Ala Gly	
959	370 375 380	
960		
961	TCG AGC AGG CTT GAT CTT TAC GGC GCG GAT TTT GGA TGG GGT AAG GCG	1199
962	Ser Ser Arg Leu Asp Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala	
963	385 390 395	
964		
965	GTG AAG CAA GAG ATA CTG TCG ATT GAT GGA GAG AAG TTT ACG ATG TCG	1247
966	Val Lys Gln Glu Ile Leu Ser Ile Asp Gly Glu Lys Phe Thr Met Ser	
967	400 405 410 415	
968		
969	TTG TGT AAA CCG AGG GAT GCT GCC GGA GGA TTG GAG GTT GGA TTG TCT	1295
970	Leu Cys Lys Pro Arg Asp Ala Ala Gly Gly Leu Glu Val Gly Leu Ser	

RAW SEQUENCE LISTING
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971		420	425	430	
972					
973	TTG CCA AAG GAG GAA TTG CAA GCT TTT GAT GAT TAT TTT GCG GAG GGA				1343
974	Leu Pro Lys Glu Glu Leu Gln Ala Phe Asp Asp Tyr Phe Ala Glu Gly				
975		435	440	445	
976					
977	ATA AAG GGT TGATTAATCA TTTAATCATG TATTATGAAG TTGGATGAAA				1392
978	Ile Lys Gly				
979		450			
980					
981	TCCTCTGTTT CATCTCTATT GTTTAAACAA TAATTTTTTTT CCATTGAACT TTTTGTGAGTC				1452
982					
983	AATAAAAAAAAA AAAAAAAAAA AAAAAAATG AAAAAACTCA GTTATTTTTT TTTTTTTTTT				1512
984					
985	TTTTTTTTTT				1521
986					
987					

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/894,356DATE: 03/25/98
TIME: 11:00:53*INPUT SET: S24372.raw*

Line	Error	Original Text
865	Wrong Nucleic Acid Designator	-TG ACC ACC CTC CTC GAA TCC TCC CGA GTG GCG



Notice of Availability of Checker Program

Applicant Aid for Biotechnology Computer Readable Form (CRF)
Sequence Listing Submissions

The Patent and Trademark Office (PTO) has developed a computer program, called Checker, that will aid applicants in identifying and correcting errors prior to making submissions for compliance with the Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures (Sequence Rules: 37CFR 1.821 through 1.825). Final rules were published in the *Federal Register* (55 FR18230) on May 1, 1990, and in the PTO *Official Gazette* (1114 Off.Gaz.PatOffice 29) on May 15, 1990.

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The software can be accessed/requested from the following locations:

- 1) Dial-up access through the Internet. Location is <ftp://ftp.uspto.gov>
The software is in current directory: `pub/checker/`
Download all the files. Cost: Free-of-charge
- 3) For diskette copies, mail to: U.S.P.T.O., OEIP, CRYSTAL PARK 3, SUITE 441
WASHINGTON DC 20231

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PTO Deposit Account

For Further Information, Contact: **Arti Shah at 703-308-4212**